

Fri Jul 2 09:20:41 2004

US-09-445-223-2.

GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: July 2, 2004, 03:26:18 ; Search time 154 Seconds  
(without alignments)  
7560.305 Million cell updates/sec

Title: US-09-445-223-2  
Perfect score: 2098  
Sequence: 1 ggccattatggatgggg.....aaaaaaaaaaaaaaaaaa 2098

Scoring table: OLIGO\_NUC  
Gapop 60.0 , Gapext 60.0

Searched: 682709 seqs, 277475446 residues

Word size : 0

Total number of hits satisfying chosen parameters: 839752

Minimum DB seq length: 0  
Maximum DB seq length: 50

Post-processing: Listing first 45 summaries

Database : Issued Patents NA.\*  
1: /cgn2\_6/ptodata/2/ina/5A\_COMB.seq:\*  
2: /cgn2\_6/ptodata/2/ina/5B\_COMB.seq:\*  
3: /cgn2\_6/ptodata/2/ina/6A\_COMB.seq:\*  
4: /cgn2\_6/ptodata/2/ina/6B\_COMB.seq:\*  
5: /cgn2\_6/ptodata/2/ina/PCTUS\_COMB.seq:\*  
6: /cgn2\_6/ptodata/2/ina/backfiles1.seq:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

RESULT 2  
US-08-113-646A-44  
; Sequence 44, Application US/08113646A  
; Patent No. 5578468  
; GENERAL INFORMATION:  
; APPLICANT: PICKUP, David J.  
; APPLICANT: PATEL, Dhaval Kumar  
; APPLICANT: ANTCHAK, James B.  
; TITLE OF INVENTION: SITE-SPECIFIC RNA CLEAVAGE  
; NUMBER OF SEQUENCES: 44  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: NIXON & VANDERHYE P.C.

